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Cassava 由来 SHNL の ^{Amino acid seq.} アミノ酸配列 (Gene Bank No. AJ223281)

MAVVDFVLIHTICHCAWIWYKLKPVLEAAGHKVLTALDLAASGVDPQRQIEQINSFDEYSEPLLTFMESLPQ
GEKVILVGESCGGLNIAAADKYPEKIAAAVFQNSLLPDTKIIKPSYVVDKLMFVFPDWKDTEYFEFSNSN
GETITGMVLGLKLMRENLYTICPPEDYELAKMLTRRGSLFQSILAQREKFTEKGYGSIKKIYVWTGDDKI
FLPEFQLWQJENYKPDVFRVMGGDIKQLTKTNEIAGILQKVADIYA

Prunus 由来 RHNL の ^{Amino acid seq.} アミノ酸配列 (Gene Bank No. AF043186)

MEKSTMSAVVLVLNLLVLHLQYSEVHSLANTSSEHDFGYLK FVYNAVDLELECSYDYIIVGGGTSGCPLA
ATLSANYSVLVLERGTIATEYPNTLTVDGFAYNLQQQDDGKTPVERFVSEDGIDNVRSLGGTTIINAGV
YARANESFYNNSGV EWDLDLVNEAYEWVEDAIVYKPSNQSWQSITGTAFL EAGVHPDNGFGLVHEEGT
RLTGSTFDNSGTRHASDELLNKGD PDLNKVAVEAAVQKII FSTESSGLTAVGVVYTDSNGTSHRALVSGK
GEVILSAGTLGTPQLLLLLSGVGPE SYLTSLNISVVASHPYVQQYVNDNPRNF INILPPNPIEPSTVTVLGITS
DFYQCSLSSLFPDTPPFSLFPTTSYPLPNQTF AHIVSKVPGPLSAGSLTLQSSSNVSVAPNVKFNYCSDPV
DLTHCVSGMKKIGVFLSTDALKPYKVDDLPGIDGFNILGTPLPENQTDDAAFEKFCRDTVASYWHYHGG
AIVGKVIDGNFRVTGINALRVVDGSTFPATPASHPQGFYLM LGRYVGTKIVQERSASGEA IHTSTFKPKLM
DSLKSALSFAF

両者のホモロジー検索結果

[GENETYX : Maximum Matching]

Date : 2002.12.17

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Sequence 1 : M.esculenta SHNL AA seq
Size : 258
Matching Position : 1 - 258

Sequence 2 : P.serotina RHNL AA seq
Size : 574

Matching Position : 1 - 574

Matching Condition.

Matches : -1
Mismatches : 1
Gaps : 1
*N+ : 2

Matching : 15.68 [%]
Weight : 456

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1 : M-----AVV -DFVL--IHT ICH----- --GAWIWKYKL KPV----- LE-----
    *      ***      *      *      *      *      *      *
1 : MEKSTMSAVV LVLNLLVLHL QYSEVHSLAN TSSEHDFGYL KFYVNAVDLE LEGSYDYIIV

28 : ----- --AA----- --GHKVT- ----- ALDLAASGVD PRQIEQINSF
    *      *      *      *      *      *
61 : GGGTSGCPLA ATLSANYSVL VLERGTIATE YPNTLTVDGF AYNLQQQDDG KTPVERFVSE

55 : D----EYSEP L----- --LTFMESLP QGEKV----I LVGESCGGLN IAI-----
    *      *      *      **      *      *      *
121 : DGIDNVRMRI LGGTTIINAG VYARANESFY NNSGVEWDLD LVNEAYEWVE DAIVYKPSNQ

89 : ----- -AADKYPEKI AAIV----- -----FQNS- -----LL- ---PD-----
    *      *      *      *      *      *
181 : SWQSITGTAF LEAGVHPDNG FGLVHBEGTR LTGSTFDNSG TRHASDELLN KGDPDNLKVA

110 : -----TKHKP SYVVDKLMEV FPDWKD----- -----TEYF EFSNSNGETI TGMVLGL---
    *      *      *      *      *      *      *
241 : VEAIVQKIIF STESSGLTAV GVYITDSNGT SHRALVSGKG EVILSAGTIG TPQLLLLSGV
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152 : -----KLMREN LYTICPPEDY E-----
      * * * * *
301 : GPESYLTSLN ISVVASHPYV GOYVNDNPRN FINILPPNPI EPSTVTVLGI TSDFYQCSLS
169 : LAKMLTRRGS LF----- --QS---IL AQREKF---
      * * * * *
361 : SLFFDTPPFS LFPTTSYPLF NQTFHIVSK VPGPLSAGSL TLQSSSNVSV APNVKFNYS
191 : -----TEKGY GSIK----- ----KIYV-- -WIGDD---- -KIFLPEFQ- -----
      * * * * *
421 : DPVDLTHCVS GMKKIGVFLS TDALKPYKVD DLPGIDGFNI LGTPLPENQT DDAAFEKFCR
217 : ----- LWQIENYKPD LVFRVMG--- -----GDH KLQL-----T
      * * * * *
481 : DTVASYWHYH GGAIVGKVID GNFRVTIGINA LRVVDGSTFP ATPASHPQGF YLMLGRYVGT
242 : K----- ---TNEIAG IL---QKVAD IYA-
      * * * * *
541 : KIVQERSASG EAIHTSTFKP KLMDSLKSAL SFAF

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[GENETYX : Amino Acid Sequence Homology Data]

Date : 2002.12.17

1st Amino Acid Sequence

File Name : Cassava SHNL AA seq.

Sequence Size : 258

2nd Amino Acid Sequence

File Name : P.serotina RHNL AA seq

Sequence Size : 574

Unit Size to Compare = 2

Pick up Location = 1

[31.818% / 22 aa] INT/OPT.Score : < 21/ 63 >

1' MAVVDFVLIIH TICHGAWIWY KLRPVLEAAG HKVTALDLAA SGVDPRQIEQ INSFDEYSEP
1" MEKS TMSAVVLVLN LLVLHLQYSE
61' LLTFMESLPQ GEKVILVGES CGGLNIAIAA DKYPEKIAAA VFQNSLLPDT KHKPSYVVDK
25" VHSLANTSSE HDFGYLKFEVY NAVDLELEGS YDYIIVGGGT SGCPLAATLS ANYSVLVLER
121' LMEVFPDWKD TEYFEFSNSN GETITGMVLG LKLMRENLYT ICPPEDYELA KMLTRRGSLE
85" GTIATEYPNT LTVDGFAYNL QQQDDGKTPV ERFVSEDGID NVRSRILGGT TIINAGVYAR
181' QSILAQREKF TEKGYGSIKK IYVWTGDDKI FLPEFQLWQI ENYKPDLVFR VMGGDHKLQL
... * * . . . * . * * *
145" ANESFYNNNG VEWDLDLVNE AYEWEIDAIV YKPSNQSWQS ITGTAFLEAG VHPDNGFGLV
241' TKTNEIAGIL QKVADIYA
205" HEEGTRLTGS TFDNSGTRHA SDELINKGDP DNLKVAVEAA VQKIIFSTES SGLTAVGVVY